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P 2ahl

Mutani Mus Musculus P38 Kinase Domain in Complex with Inhibitor PG-874743



Release Date: 11-Apr-2006 Exp. Method: Characteristics Resolution: 2.10 Å

Transferase

Compound

Polymer: 1 Molecule: Mitogen-activated kinase 14 Mutation: Y182F, T180A C

no.: 2.7.1.37 ES

Authors

Walter, R.L., Mekel, M.J., Evdokimo A.G., Pokross, M.E., Brugel, T.A.

₹ 2ahm

Mutated MAP kinase P38 (Mus Musculus) in complex with Inhbitor PG-895449

Characteristics

Release Date: 02-May-2006 Exp Method: Resolution: 2.35 Å

Transferase Classification

Compound

Polymer: 1 Molecule: Mitogen-activated 14 Mutation: Y182F, T180A Chains A

no.: 2.7.1.37 🗐

Walter, R.L., Mekel, M.J., Evdokimo Authors A.G., Pokross, M.E., Maier, J.A.

Mutated Mouse P38 MAP Kinase

₹ 2gim

n e Domain in complex with inhibitor PG-892579

Characteristics

Release Date: 23-May-2006 Exp. Method:

Besolution: 1.90 Å Transferase Classification

Compound

Polymer: 1 Molecule: Mitogen-activated

14 Chains: A FC no.: 2.7.11.24 89



http://www.rcsb.org/pdb/results/results.do?gotopage=4

Walter, R.L., Mekel, M.J., Evdokimov. Authors A.G., Pokross, M.E., Sabat. M.

Mutated MAP kingse P38 / Mus Musculus) in complex with Inhbitor

PG-951717

Release Date: 11-Jul-2006 Exp. Method: Characteristics Resolution: 1.80 Å

Classification Transferase

Polymer: 1 Molecule: Mitogen-activated Compound

kinase 14 Chains A EC no : 2.7.11.24 Walter, R.L., Makei, M.J., Evdokimo

Authors A.G., Pokross, M.E., Sabat, M.

Structural studies of profein P 2502 tyrosine phosphalase bela catalytic domain in complex with inhibitors

> Release Date: 13-Jun-2006 Exp. Method: Characteristics

Resolution: 2.30 Å

Hydrolase Classification

Polymer: 1 Molecule: Protein tyrosine p receptor type, B, Fragment: catalytic d Compound residues 1662-1973 Chains: A,B EC

no.: 3.1.3.48 🧐

Evdokimov, A.G., Pokross, M.E., W R.L., Mekel, M., Gray, J.L., Peters, M.B., Amarasinghe, K.D., Clark, C.P. Authors

Structural studies of protein

tyrosine phosphatase beta cafalytic domain in complex with inhibitors

Hydrolase

Classification

Classification

Release Date: 13-Jun-2006 Exp. Method: Characteristics

Resolution: 1.65 Å

Polymer: 1 Molecule: Protein tyrosine p

receptor type, B, Fragment: catalytic d 1970 Mutation P1748G, C1749G Ch Compound

no.: 3.1.3.48 🧐

Evdokimov, A.G., Pokross, M.E., W.R.L., Mekel, M., Gray, J.L., Peters, M.B., Amarasinghe, K.D., Clark, C.N. Authors

Structural studies of protein ₹ 2h04 tyrosine phosphatase bela catalytic

Hydrolase

domain in complex with inhibitors Release Date: 13-Jun-2006 Exp. Method:

Characteristics Resolution: 2.30 Å

Polymer: 1 Molecule: Protein tyrosine p

receptor type, B, Fragment: catalytic d













































residues 1662-1973 Chains: A EC no. 3.1.3.48 59 Compound

Evdokimov, A.G., Pokross, M.E., Waiter, R.L., Mekel, M., Gray, J.L., Peters, K.G., Maier, M.B., Amarasinghe, K.D., Clark, C.M., Nichols, Authors

Grystal structure of HIF proly! ₹ 2hbt hydroxylase EGLN-1 in complex with a biologically active inhibitor

Release Date: 27-Jun-2006 Exp Method: Characteristics

Resolution: 1.60 Å Classification Oxidoreductase

Polymer: 1 Molecule: Egl nine homolog

Compound Fragment: catalytic domain Chains: A no.: 1.14.11.-

Evdokimov, A.G., Walter, R.L., Mek M., Pokross, M.E., Kawamoto, R., E Authors

Crystal structure of HIF prolyt ₽ 2hbu

Classification

P 2hc1

nydroxylase EGLN-1 in complex with a biologically active inhibitor Release Date: 27-Jun-2006 Exp. Method.

Characteristics Resolution: 1.85 Å Oxidoreductase

Polymer: 1 Molecule: Egl nine homolog Compound Fragment: catalytic domain Ghains: A

по.: 1.14.11.-

Evdokimov, A.G., Walter, R.L., Mek M., Pokross, M.E., Kawamoto, R., E Authors

Engineered catalytic domain of protein tyrosine phosphatase HPTPbeta.

Release Date: 27-Jun-2006 Exp Method: Characteristics

Resolution: 1.30 Å Hydrolase

Classification

Polymer: 1 Molecule: Receptor-type tyre Compound phosphatase beta Fragment: catalytic

Mutation: ves Chains: A EC no.: 3.1.3.4 Evdokimov, A.G., Pokross, M., Wall

Authors A., Mekel, M.

(123456C)

@ RCSB Protein Data Bont